AMENDMENTS TO THE SPECIFICATION

Please replace the paragraph starting on page 15, line 1 with the following amended paragraph:

The amino acid sequence of the purified material was determined using a model 477A sequencer (Applied Biosystems, Foster City, USA) and found to be: Met-Ala-Gly-Asn-Ser-Ser-Asn-Phe-Ile-His-Lys-Ile-Lys-Gln-Ile-Phe-Thr-His-Arg (19 residues~~)~~, ~~(seq. id. No. 1~~residues 19-37 of SEQ. ID. NO: 3) with a calculated molecular weight of 2229.6. This amino acid sequence corresponds with the amino acid sequence encoded by part the DNA sequence that was determined for DNA from *Lactobacillus sake* LTH673 (Fig. 2).

Please replace the paragraph starting on page 15, line 32 to page 16, line 2,with the following amended paragraph:

Experiments similar to the ones described in examples 1 and 2 were conducted with the bacteriocin producing strain *Lactobacillus plantarum* C11. These experiments yielded results that were essentially similar to the results described in examples 1 and 2. The sequence of the amphiphilic inducing peptide for the *Lactobacillus plantarum* C11 regulatory system is: Lys-Ser-Ser-Ala-Tyr-Ser-Leu-Gln-Met-Gly-Ala-Thr-Ala-Ile-Lys-Gln-Val-Lys-Lys-Leu-Phe-Lys-Lys-Trp-Gly-Trp (26 residues) (~~seq. id. No. 2~~SEQ. ID. NO: 1).

IN THE SEQUENCE LISTING

Please substitute the Sequence Listing attached hereto for the Sequence Listing filed February 1, 1999 and presently of record.